

IN THE CLAIMS

Please amend the claims as indicated in the following listing of claims, which replaces all previous listings of claims.

1-44. (Cancelled)

45. (Previously presented) A nucleic acid vector comprising sequences encoding *E. coli* argU, ileY and leuW tRNAs, wherein the backbone of said vector is derived from pACYC 184.

46. (Previously presented) The vector of claim 45 wherein said sequences encoding said *E. coli* tRNAs are present in the order argU, ileY, leuW.

47. (Previously presented) The vector of claim 46 wherein said sequence encoding ileY tRNA is present in the opposite orientation to said sequences encoding argU and leuW tRNAs.

48. (Previously presented) The vector of claim 45 that comprises a *tet* promoter.

49. (Previously presented) The vector of claim 48 wherein said *tet* promoter is operably linked to a sequence encoding a said tRNA.

50. (Previously presented) The nucleic acid vector of claim 45 wherein said sequence encoding *E. coli* argU tRNA comprises the sequence between base pairs 8041 and 8260 of GenBank Accession No. AE000159.

51. (Previously presented) The nucleic acid vector of claim 45 wherein said sequence encoding *E. coli* ileY tRNA comprises the sequence between base pairs 7741 and 7950 of GenBank Accession No. AE000350.

52. (Previously presented) The nucleic acid vector of claim 45 wherein said sequence encoding *E. coli* leuW tRNA comprises the sequence between base pairs 241 and 378 of GenBank Accession No. J01713.

53. (Previously presented) The nucleic acid vector of claim 45 which further comprises a

chloramphenicol resistance gene.

54. (Currently amended) ~~A~~ An isolated nucleic acid comprising a promoter operably linked to sequences encoding *E. coli* argU, ileY and leuW tRNAs.

55. (Previously presented) The nucleic acid of claim 54 wherein said sequences encoding *E. coli* tRNAs are present in the order argU, ileY, leuW.

56. (Previously presented) The nucleic acid of claim 55 wherein said sequence encoding ileY tRNA is present in the opposite orientation to said sequences encoding argU and leuW tRNAs.

57. (Previously presented) The nucleic acid of claim 54 wherein said promoter is the *tet* promoter.

58. (Previously presented) The nucleic acid of claim 54 wherein said promoter is endogenous to a said sequence encoding an *E. coli* tRNA.

59. (Previously presented) The nucleic acid of claim 54 wherein said sequence encoding *E. coli* argU tRNA comprises the sequence between base pairs 8041 and 8260 of GenBank Accession No. AE000159.

60. (Previously presented) The nucleic acid of claim 54 wherein said sequence encoding *E. coli* ileY tRNA comprises the sequence between base pairs 7741 and 7950 of GenBank Accession No. AE000350.

61. (Previously presented) The nucleic acid vector of claim 54 wherein said sequence encoding *E. coli* leuW tRNA comprises the sequence between base pairs 241 and 378 of GenBank Accession No. J01713.